

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN	435	6

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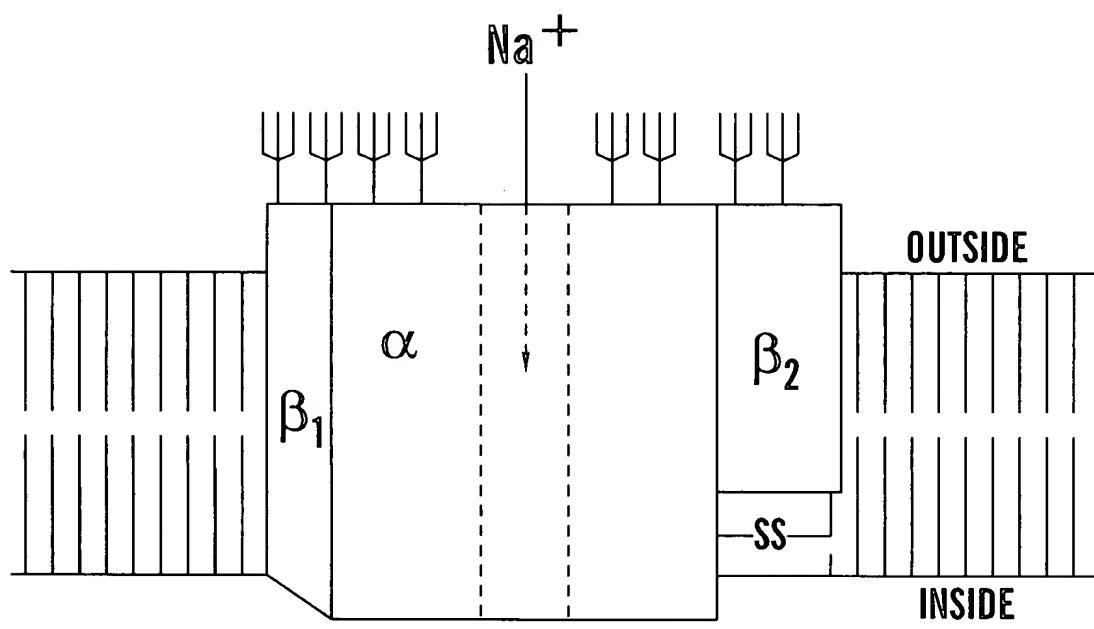
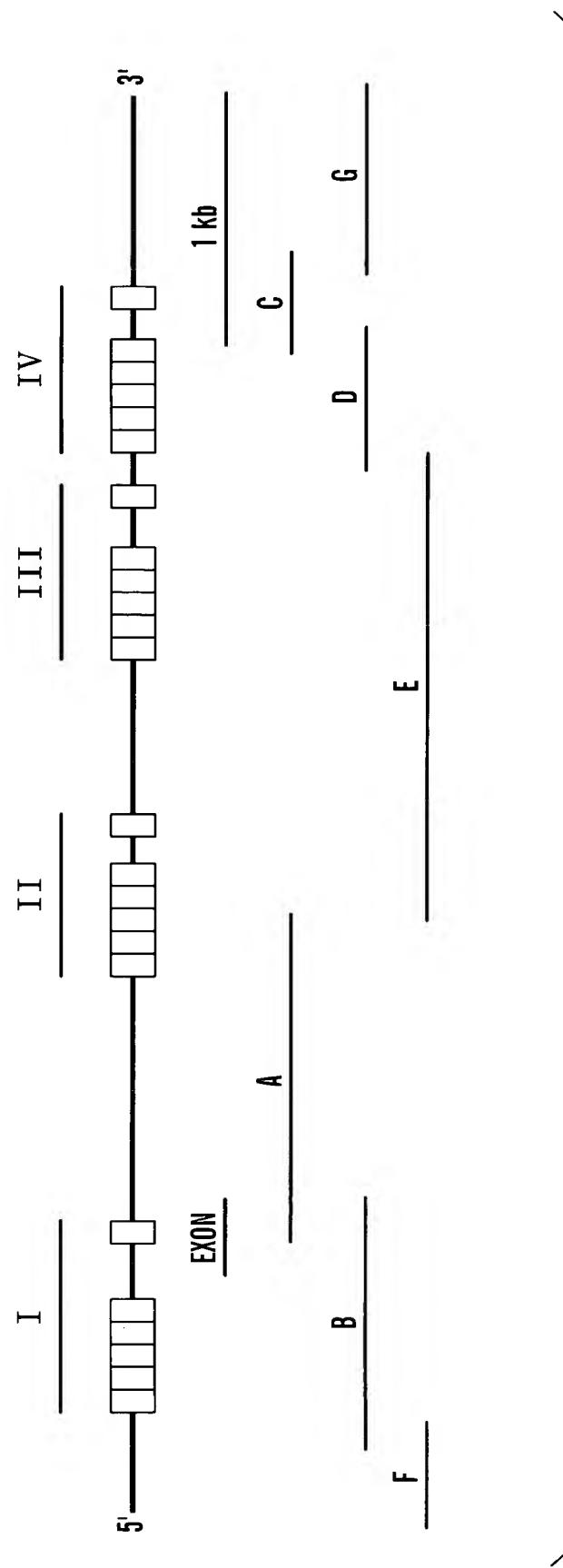


FIG. 1

65-63371-102692

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**FIG. 2**

APPROVED BY RAFTSMAN	O.G. FIG.	
	CLASS 436	SUBCLASS 6

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Nайд. МТЕДСДСИСЕЕРSLRPFTRSLLQIEQRIA. EHEKQELEKRAEGERE..... QIRYDDEDEGPQDPPTLEQGVPIPVRMQ 79  
 538ge ----- L ----- A ----- Q ----- M- 79  
 para ----- V ----- A ----- . ----- VPRYGRKKKQKE ----- L- 90  
IS1 -----  
 GSFPELASTPLIEDIDPFSNVLTFTVVISKGKDIFRFSASKAMWLDPFNPIRRVAIYLVHPLFSLFIFTILTNCIMIMPTPTVESTEVIFTGIYT 179  
 ----- F ----- I ----- L ----- T ----- 179  
 ----- Y ----- V ----- M ----- V ----- 190  
IS2 -----  
IS3 -----  
 FESAVKVMARGEFILECPFTYLRDAWNWLDFTVIALAYTMGIDLGNLAALRTRVLRAKTVAIIVPGLKTTIVGAVIESVNLRDVILTMFSLSVFALMGL 279  
 ----- ----- ----- ----- 279  
 ----- ----- ----- ----- 290  
IS4 -----  
 QIYMGVLTQKCIKREPLDGSWGNLTDENWFLHNSNSMWTENDGESYPVCGENVSGAGQCGGEDYVCLQGFGPNPNYDYTTSFDSEFGWAELSAFRMLMTQDFW 379  
 ----- Q ----- R ----- FL ----- S ----- FT-ND-E-Y-V ----- V ----- GE ----- D ----- 379  
 ----- E ----- K ----- DY ----- R ----- YS-DE-I-F-L ----- I ----- DD ----- G ----- 390  
IS5 -----  
 EDLYQHVLQAAGPWHMLFIFTIFLGSFYLVNLILATVAMSYDELQKKAEEEEEAAIREAEEAAAAKAKLEERANVAQQAAQDAADAAAAALHPEM 479  
 ----- H- Q ----- K ----- VA ----- Q ----- D-AA ----- 479  
 ----- L- R ----- R ----- AQ ----- A ----- A-EE ----- 490  
 \* -----  
 AKSPTYSCISYELFVGGEKGNDDNNKERSIRSVESESVSVIQRQPAPTTAP. ATKVRKVSTTISLPGSPFNLLRGSRSSHKYTIRNGRGRFGIPGS 578  
 ----- P ----- L ----- 578  
 ----- HQ ----- I ----- 590

FIG. 3A

APPROVED	O.G. FIG.	
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DRKPLVLTQTYQDAQQHLPLYADDSSNAVTTPMSEENGAIITVPAVYCNIGSRHSSYTSHQRISYTSHGDLJGGMAAMGASTMTKESKLRSRNTRNQSIGAATN 678  
 Q-----A-C-----A-A-----S-----I-ATN 678  
 S-----V-G-----V-V-----N-----V-TNG 690

GGSSTAGGGYPDANHK. EQRDYEMGQDYTDEAGKIKHDMNPFIEPVQTQTVDMKDVINDIIEQAAGRHSRASERGEDDEDGPTFKDIALEYILKGI 777  
 -GSSTAGGGYP-A---EQ---M-QDY-----E-----I-Y-----777  
 -TTCL. . . . . -T---LDH---I-LEC-----D-----K-V-----784

||S1  
 EIFCWIDCCWVWLKFQEWVSFTIVDPFVELFTILCIVWNTMFAMDDHDMNPELEKVLKSGNYFFTATFAIEASMKLMAASPKYFQEGWNIFDFIIVAL 877  
 EI-----F-----P-L-K-----S-----877

DV-----L-----K-M-R-----T-----884

||S2  
 SILELGLLEGVQGLSVLRSFRLLRVFKLAKSMPTINLLISIMGRTMGALGNLTFLCIIIFIAVMGMQLFGKNTYDHKDRFKDHELPRWNFTDFMHSFMI 977  
 EI-----K-HE-----I-----K-HE-----977  
 -H-----P-GD-----H-----P-GD-----984

||S3  
 VFRVLGEWIESMWDCMYVGDVSCIPFFLATVIVNLFALLLSNFGSSLSAPTAADNDTNKIAEAFNRIARFKNNVKRNIADCFCKLIRNKLTNQ 1077  
 F-----A-N-----A-N-----1077

||S4  
 ||S5  
 SILELGLLEGVQGLSVLRSFRLLRVFKLAKSMPTINLLISIMGRTMGALGNLTFLCIIIFIAVMGMQLFGKNTYDHKDRFKDHELPRWNFTDFMHSFMI 977  
 EI-----K-HE-----I-----K-HE-----977  
 -H-----P-GD-----H-----P-GD-----984

||P  
 ||S6 ◇  
 ISDQPSEHGDNEELGHDEIMGDGLIKKGMKGETQLEVAIGDGMEFTIHDMNNKPKKSFMNTTMIGNSINHQDNRLHEHNRGLSIQDDDTASIN 1177  
 MG-----M-GE-----FI---T-MIGNSINHQDNRLHEHNRGLSIQ-----1177  
 LA-----I-EQ-----YL-A-----1159

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FIG. 3B

APPROVED	O.G. FIG.	
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 °  
 SYGSHKJNRPFKDESHKGSAETTIEGEEKRDKVSKEDLGLDEELDEAEGDEGQLDGDDIIHQNDDEIIDDYPADCFPDSYYKKFPILAGDEDSPFWQGWGN 1277  
 I-----V-----A-GD-Q-----Q-N-DE-I-D-----F-----E-----1277  
 M-----A-----G-CE-P-----H-ED-L-E-----C-----D-----1258  
III S 1  
 LRLKTFQLIENKYFETAVITMILMSSIALAEDVHLPRPVMQDILYMDRIFTVFFLEMILKWMALGFKVYFTNAMCWLDFTVIMLSLNLVAVWSGL 1377  
 Q---N---D---VM---F---L---L---VWS-L 1377  
 R---D---Q---IL---L---V---F---SLV-A 1358  
III S 2  
 #  
 NDIAVFRSMRTLRAVRPLRAVSRMKGKVVNNALVQAIPSIINVLLVCLIFWLIFAMGVQLFAGKYFKCKDGNNDTLSHEIIIPRNACKSENITWENSA 1477  
 ND-AV-RS---V--WE--K-----K-G-D-V-----E-----1477  
 GG-QA-KT---M---MQ---R-----E-M-G-K-----E-----V---1458  
III P  
 MNFDHGNAYLCLFQVATFKGWIQIMNDAIDSREVDKQPIRETNIYMYLYFVFLIFGSFTTINLFIGVITIDNNEQKKKAGGSLEMFTEDQKKYNNAM 1577  
 °  
 KKMGSKKPLKAIPPRPRWRPQAIVFEIVTDDKKFDIIIMLFIGLNMFMTLDRYDAEAYNNVLDKLNGIFVVFISGECLLKTFALRYHYFKEPWNLFDVVV 1677  
 EA---N---K---G-----K-----G-----1677  
 DT---A---Y---A-----I-----1658

FIG. 3C

APPROVED	O.G. FIG.	
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**IVS4**

VILSILGLVLSDIEKYFVSPTLLRVRVAKVGRVLRLVKGAKGIRTLFALAMSLPNCILLLFLVMFIFAIIFGMSFFMIVKEKSGTINAVYNFKTFG 1777

-D-----A-----1777

-D-----D-----1758

**IVP**

QSMILLFQMTSAGMDGVLDAAIINEEDCDPPNDKGYPGNCGSATVGITFLLSYLVISFLIVINMYTAVILENYSQATEDVQEGLTDDDDMYYTEIWQQF 1877

-D-----A-----1877

-A-----A-----1858

DPEGTQYTRYDQLSEELDVEPPLQIHKPNKYKTIISMDMPICRGDMYCVDILLDALTKDFEARKGNPPIEETGEIGEIAARPDTEGYDPPSSTLWROREEX 1977

-M-----M-----1977

-I-----L-----1958

◆◆◆◆◆◆◆◆◆◆  
CAKLQNAWRRYK.....NGPPQEGDEGEAAGGEDGAEGGEGGGCGSGGGGDDGGSATGATAAGAT...SPSPDPDAGEADGASVG...GPLSPGCV 2063  
-K---N---RY-.....NGPPQE--E-EAAG-EDGAEKGEGGGSG--GDDDG-S-T....-AGAT..SPTDPD-GE-DG-SAGNGG-PLSP-CV 2062  
-R---H---KH-ARGE GGGSFEPDTDH--G-DPDA-DPAPDEATDGA PA--DGSVN-T-E--AD-DESNVNNSPGEDA A A-A-A-TTTA-SP 2055

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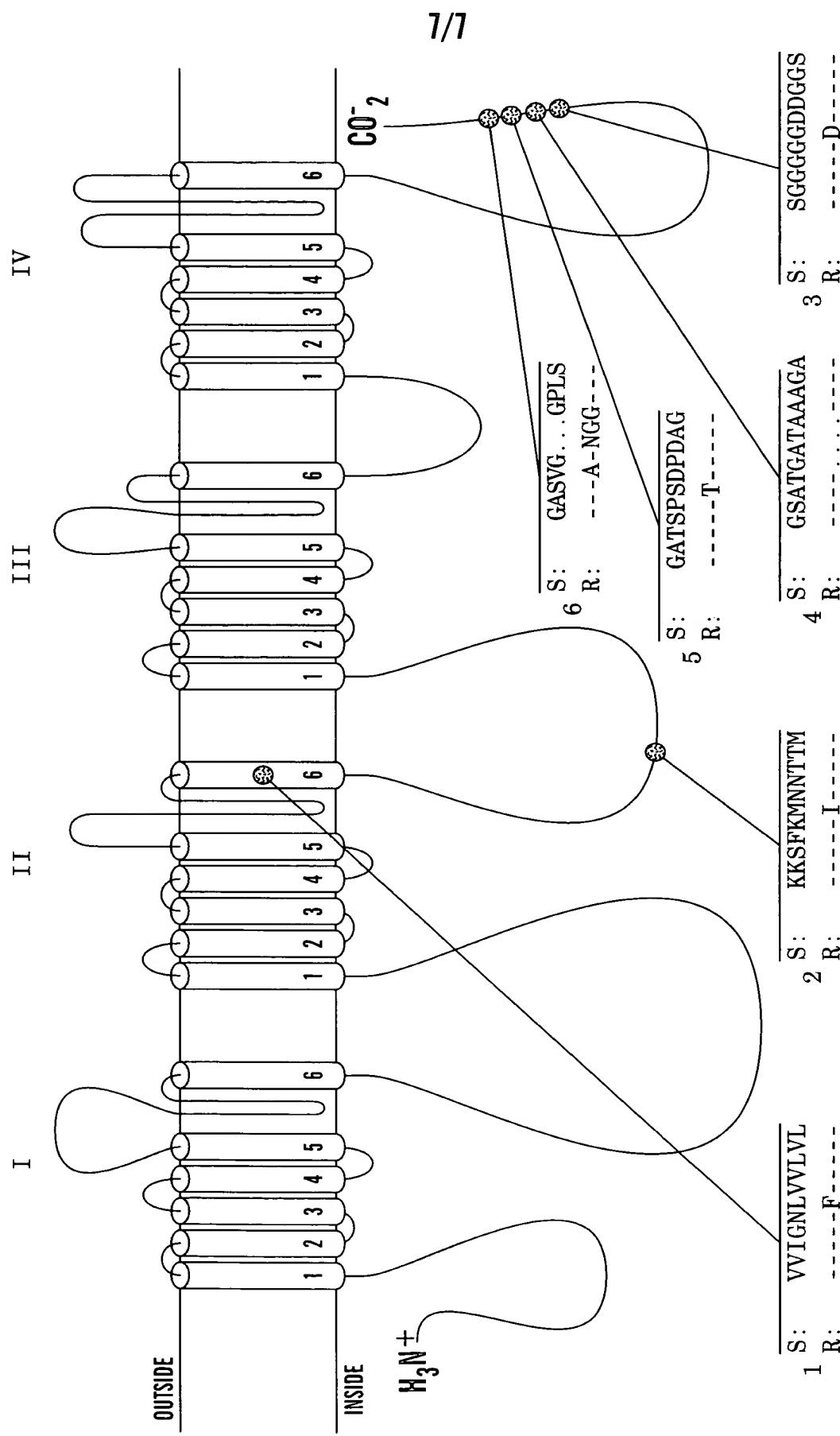
SGGSNNGRQTAVLVESDGFTVTKNGHKVTHSRSPSITSRTADV 2105

SG--N-----2104

GA--A-----2100

**FIG. 3D**

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